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Sea Sector

Docket No.: PA-0033 US

I hereby certify has this correspondence is being deposited with the United States Postar Friends at class mail in an envelope addressed to: Commissioner for Patents, Attn: Box Missing Parts Washington, D.C. 18231 of Page 18, 2001.

Printed name:

Katherine Stotes

#### IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of: Schebye et al.

Title:

CDNAs EXPRESSED IN ADIPOCYTE DIFFERENTIATION

Serial No.:

09/918,624

Filing Date:

7/30/01

Examiner:

Duffy, P.

Group Art Unit: 1645

Commissioner for Patents Attn: Box Missing Parts

Washington, D.C. 20231

# RESPONSE TO NOTICE TO FILE MISSING PARTS, FILING DATE GRANTED AND NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Sir:

In response to the Notice to File Missing Parts of Application, Filing Date Granted and Notice to Comply with Requirements for Patent Applications Containing Nucleotide Sequence and/or Amino Acid Sequence Disclosures mailed by the United States Patent and Trademark Office on October 18, 2001, Applicants submit the following documents to complete the filing for the above-identified patent application:

- 1. Copy of Notice to File Missing Parts of Application, Filing Date Granted (2 pp.);
- 2. Executed Declaration and Power of Attorney for United States Patent Application (6 pp., in counterpart);
- 3. Copy of Notice to Comply with Requirements for Patent Applications Containing Nucleotide Sequence and/or Amino Acid Sequence Disclosures (4 pp.);
  - 4. Submission Under 37 CFR § 1.821-1.825 Sequence Listing (1 pg.);
  - 5. Compact Disk-Recordable (CD-R) discs containing Sequence Listing(3 CD-Rs); and
  - 6. Return postcard.

Docket No.: PA-0033 US

Please charge the following fees to **Deposit Account No. 09-0108**:

Basic surcharge fee for filing the declaration:

130.00

Total fees charged to Deposit Account No. 09-0108:

130.00

If the Patent Office determines that additional fees are due or that an excess fee has been paid, the Patent Office is authorized to debit or credit (respectively) **Deposit Account No. 09-0108.** 

If there are any questions regarding the above, the Examiner is invited to call the undersigned at 650-855-0555.

This response is enclosed in duplicate.

Respectfully submitted,

INCYTE GENOMICS, INC.

Date: 18 Docume 2001

Lynn E. Murry, Ph.D.

Reg. No. 42,918

Direct Dial Telephone: (650) 845-4159

3160 Porter Drive Palo Alto, CA 94304 Tel: 650-855-0555

Fax: 650-849-8886

I hereby certify that this correspon s being deposited irst class mail in with the United States Postal Servi an envelope addressed to: Commissioner for Patents, Washington, D. 2023 on December 18, 2001

Docket No.: PA-0033 US

Printed: Katherine Stofer

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

ants:

Xiao Min Schebye, Thierry R. Sornasse

CDNAS EXPRESSED IN ADIPOCYTE DIFFERENTIATION

Serial No.:

09/918,624

Filed:

July 30, 2001

Examiner:

Duffy, P.

Group Art Unit:

1645

Commissioner for Patents

Box Sequence

Washington, D.C. 20231

#### SUBMISSION UNDER 37 CFR §1.821- 1.825 SEQUENCE LISTING

Sir:

1. In accordance with the requirements of 37 CFR §1.821- 1.825, and by waiver of 37 CFR § 1.2 and in compliance with new rule 37 CFR §1.52(e), Applicants hereby submit (in duplicate) two (2) identical Compact Disk-Recordable (CD-R) discs, each containing the Sequence Listing for the above-identified application. The following files are contained on the compact discs:

File name

Size (Kbytes)

Creation Date Identifying Information

e.g.

pa0033.txt

206KB

December 18, 2001

Sequence Listing, SEQ ID NO:1-71.

These CD-Rs are provided in compliance with 37 CFR §1.821 et seq. and as permitted by 37 CFR §1.52(e). Each file recorded on the CD-R is formatted in plain ASCII text, (Sequence Listing), to be viewed on an IBM Compatible desktop computer. The files are readable using any standard word processing (Sequence Listing) (e.g., MS Word '97 or MS-Wordpad for the Sequence Listing.

2. Additionally, Applicants herein submit one (1) CD-R containing the computer readable form (the CRF) of the Sequence Listing of the above-identified application, in accordance with 37 CFR §1.821(e) and as specified by 37 CFR §1.824(a)(2).

File name pa0033.txt

Size (Kbytes) Creation Date Identifying Information

e.g.

206KB

December 18, 2001

Sequence Listing, SEQ ID NO:1-71.

KgNo. 37,027

3. The content of the Sequence Listing recorded on the two (2) CD-Rs, referred to in paragraph 1. above, is identical to the Sequence Listing presented in the CRF, referred to in paragraph 2. above, as required under 37 CFR §1.821(f).

Date: 8 Peramber 2001

Respectfully submitted,

CYTE GENOMICS, INC.

Lynn E. Murry, Ph.D.

Reg. No. 42,918

Direct Dial Telephone: (650) 845-4159

3160 Porter Drive Palo Alto, California, 94304 Tel. No. 650-855-0555 Fax. No. 650-849-8886

89190

09/918,624

1





## United States Patent and Trademark Office

COMMISSIONER FOR PATENTS UNITED STATES PATENT AND TRADEMARK OFFICE WASHINGTON, D.C. 20231

www.uspto.gov

PPLICATION NUMBER

GENOMICS, INC.

FILING/RECEIPT DATE

FIRST NAMED APPLICANT

ATTORNEY DOCKET NUMBER

09/918.624

orter Drive Palo Alto, CA 94304 07/30/2001

Xiao Min Schebye

PA-0033 US

**CONFIRMATION NO. 3179** 

**FORMALITIES LETTER** 

\*OC000000006927366\*

Date Mailed: 10/18/2001

NOTICE TO FILE MISSING PARTS OF NONPROVISIONAL APPLICATION

FILED UNDER 37 CFR 1.53(b)

Filing Date Granted

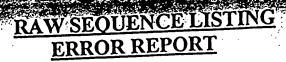
An application number and filing date have been accorded to this application. The item(s) indicated below, however, are missing. Applicant is given TWO MONTHS from the date of this Notice within which to file all required items and pay any fees required below to avoid abandonment. Extensions of time may be obtained by filing a petition accompanied by the extension fee under the provisions of 37 CFR 1.136(a).

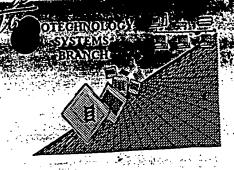
- The oath or declaration is unsigned.
- To avoid abandonment, a late filing fee or oath or declaration surcharge as set forth in 37 CFR 1.16(I) of \$130 for a non-small entity, must be submitted with the missing items identified in this letter.
- The balance due by applicant is \$ 130.
- A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing." Applicant must provide a substitute computer readable form (CRF) copy of the "Sequence Listing" and a statement that the content of the sequence listing information recorded in computer readable form is identical to the written (on paper or compact disc) sequence listing and, where applicable, includes no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b), or 1.825(d).

For questions regarding compliance to these requirements, please contact: 09918624 ■ For Rules Interpretation, call (703) 308-4216 ■ To Purchase Patentin Software, call (703) 306-2600 00000112 090108 ■ For Patentin Software Program Help, call (703) 306-4119 or e-mail at patin21help@uspto.gov or patin3help@uspto.gov 130 00 01/08/2002 SSESHE1 A copy of this notice MUST be returned with the reply.



Customer Service Center
Initial Patent Examination Division (703) 308-1202
PART 2 - COPY TO BE RETURNED WITH RESPONSE







The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: Source: Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS. PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

### Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST 25

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2Kcompliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

OIPE

RAW SEQUENCE LISTING

DATE: 08/09/2001

PATENT APPLICATION: US/09/918,624

TIME: 14:01:32

Input Set : A:\pa0033us\_seqlisting.txt Output Set: N:\CRF3\08092001\1918624.raw

**Does Not Comply** Corrected Diskette Needed

```
2 <110> APPLICANT: Schebye, Xiao Min
              Sornasse, Thierry R.
      5 <120> TITLE OF INVENTION: CDNAS EXPRESSED IN ADIPOCYTE DIFFERENTIATION
      7 <130> FILE REFERENCE: PA-0033 US
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/918,624
C--> 10 <141> CURRENT FILING DATE: 2001-07-30
     12 <150> PRIOR APPLICATION NUMBER: 60/222,470
     13 <151> PRIOR FILING DATE: 2000-07-28
     15 <160> NUMBER OF SEQ ID NOS: 71
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#### ERRORED SEQUENCES

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My blesent in the 25 house (12) port.
2191 <210> SEQ ID NO: 42
2192 <211> LENGTH: 1638
2193 <212> TYPE: DNA
2194 <213> ORGANISM: Homo sapiens
2196 <220> FEATURE:
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2197 <221> NAME/KEY: misc\_feature

16 <170> SOFTWARE: PERL Program

2198 <223> OTHER INFORMATION: Incyte ID No: g452054

2200 <400 SEQUENCE: 42 E--> 2201 (laagettagg aagatttett gggcacggta tatecagttg getaataaga aaatacgtet 60 E--> 2202 cccttcagcc tgtgccttga ctacttaaag gataggaggg aaggggagac gaagttactc 120 E--> 2203 tecteattgt gttcaccetg etecgaagaa etetgtette caetggeece tecaceteet 180 E--> 2204 ccccattete ggtagececa geetgteece ettgeecett tettacatte eggggggagg 240 E--> 2205 agggcgctgt tcagagggga ggagggcgct gttcagggag cgaaggggag cccccttgtg 300 E--> 2206 totagaaggo etetececae ecceaeceg tgtgagtttg tactgcaaag eteettggca 360 E--> 2207 tecttgeetg agttgggtgt tgggaagete aaattgeage tacaaactgg etggeageea 420 E--> 2208 ggggccggct atttaaaagc gcctgctctc ccggagcccc gtagtctctt tggaaacttc 480 E--> 2209 tgcaggggaa aagagctagg aaagagctgc aaagcagtgt gggctttttc ccttttttgc 540 E--> 2210 teetttteat tacceteet cegtttteae cetteteegg acttegegta gaacetgega 600 E--> 2211 atttcgaaga ggaggtggca aagtgggaga aaagaggtgt tagggtttgg ggtttttttg 660 E--> 2212 tttttgtttt tytttttaa tttcttgatt tcaacatttt ctcccaccct ctcggctgca 720 E--> 2213 gccaacgcct cttacctgtt ctgcggcgcc gcgcaccgct ggcagctgag ggttagaaag 780 E--> 2214 cggggtgtat tttagatttt aagcaaaaat tttaaagata aatccatttt tctctcccac 840 E--> 2215 ccccaacgcc atetecactg catecgatet cattattteg gtggttgett gggggtgaac 900 E--> 2216 aattttgtgg ctttttttcc cctataattc tgacccgctc aggcttgagg gtttctccgg 960 E--> 2217 cctccgctac ctgcgtgcac ctggcgctgc cctgcttccc ccaacctgtt gcaaggcttt 1020 E--> 2218 aattettgea actgggacet getegeagge acceeagece tecacetete tetacatttt 1080 E--> 2219 tgcaagtgtc tgggggaggg cacctgctct acctgccaga aattttaaaa caaaaacaaa 1140 E--> 2220 aacaaaaaa totooggggg coctottggc coctttatcc otgcactotc gototootgc 1200 E--> 2221 cccaccccga ggtaaagggg gcgactaaga gaagatggtg ttgctcaccg cggtcctcct 1260 E--> 2222 getgetggee geetatgegg ggeeggeeca gageetggge teettegtge actgegagee 1320 E--> 2223 ctgcgacgag aaagccctct ccatgtgccc ccccagcccc ctgggctgcg agctggtcaa 1380 E--> 2224 ggagccgggc tgcggctgct gcatgacctg cgccctggcc gaggggcagt cgtgcggcgt 1440 E--> 2225 ctacaccgag cgctgcgccc aggggctgcg ctgcctcccc cggcaggacg aggagaagcc 1500





RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/918,624

DATE: 08/09/2001 TIME: 14:01:33

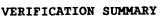
Input Set : A:\pa0033us\_seqlisting.txt
Output Set: N:\CRF3\08092001\I918624.raw

E--> 2226 gctgcacgcc ctgctgcacg gccgcggggt ttgcctcaac gaaaagagct accgcgagca 1560 E--> 2227 agtcaagatc ggtgagcgcg ctcagtgtgc cagtcagtta cgcggcgcac gggcggggg 1620 E--> 2228 cacgagaccg gctgggcc 1638



Use of n and/or Xaa has been detected in the Sequence Listing. Redow the Sequence Listing to insure a concepending explanation is presented in the <220> to <223> fields of ruch sequence using n or Xaa.





PATENT APPLICATION: US/09/918,624

DATE: 08/09/2001 TIME: 14:01:34

Input Set : A:\pa0033us\_seqlisting.txt
Output Set: N:\CRF3\08092001\I918624.raw

```
L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:339 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:346 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:355 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:356\ M:341\ W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:406 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:407 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:576 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:765 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:766 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:767 \ M:341 \ W: \ (46) \ "n" \ or "Xaa" \ used, for SEQ ID#:11
L:768 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:864 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:880 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:1267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:1268 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:1786 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:1793 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:2138 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40
L:2185 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41
L:2201 M:254 E: No. of Bases conflict, LENGTH:Input:60 Counted:59 SEQ:42
L:2201 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
M:254 Repeated in SeqNo=42
L:2228 M:252 E: No. of Seq. differs, <211>LENGTH:Input:1638 Found:1637 SEQ:42
L:2253 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43
L:2254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43
L:2255 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43
L:2338 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:2339 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:2755 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50
L:2790 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:2791 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:2803 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:2810 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:2842 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:2845 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:2988 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
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L:3193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60
L:3196 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60
L:3227 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61
L:3258 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62
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